RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: $\frac{10/537,280}{PCT-EFS}$ Date Processed by STIC: $\frac{1-9-06}{1}$

ENTERED



PCT

RAW SEQUENCE LISTING DATE: 01/09/2006 PATENT APPLICATION: US/10/537,280 TIME: 08:29:26

Input Set : N:\efs\10537280_efs\URQUP16_seq.txt

Output Set: N:\CRF4\01092006\J537280.raw

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3 <110> APPLICANT: Sanders, Jane
              Furmaniak, Jadwiga
              Smith, Bernard Rees
      7 <120> TITLE OF INVENTION: Binding Partners for the Thyrotropin Receptor and uses
thereof
      9 <130> FILE REFERENCE: URQU.P-016
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/537,280
C--> 13 <141> CURRENT FILING DATE: 2005-05-27
     15 <150> PRIOR APPLICATION NUMBER: PCT/GB2003/005171
     16 <151> PRIOR FILING DATE: 2003-11-28
     19 <150> PRIOR APPLICATION NUMBER: GB 0227964.4
     20 <151> PRIOR FILING DATE: 2002-11-29
     22 <150> PRIOR APPLICATION NUMBER: GB 0302140.9
     23 <151> PRIOR FILING DATE: 2003-01-29
     25 <150> PRIOR APPLICATION NUMBER: GB 0315147.9
     26 <151> PRIOR FILING DATE: 2003-06-27
     28 <160> NUMBER OF SEQ ID NOS: 38
     30 <170> SOFTWARE: PatentIn version 3.1
     32 <210> SEQ ID NO: 1
     33 <211> LENGTH: 121
     34 <212> TYPE: PRT
     35 <213> ORGANISM: Homo sapiens
     37 <400> SEOUENCE: 1
     39 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu
     40 1
                        5
                                             10
    43 Ser Leu Lys Ile Ser Cys Arg Gly Ser Gly Tyr Arg Phe Thr Ser Tyr
     47 Trp Ile Asn Trp Val Arg Gln Leu Pro Gly Lys Gly Leu Glu Trp Met
     48
                35
     51 Gly Arg Ile Asp Pro Thr Asp Ser Tyr Thr Asn Tyr Ser Pro Ser Phe
            50
                                55
                                                     60
     55 Lys Gly His Val Thr Val Ser Ala Asp Lys Ser Ile Asn Thr Ala Tyr
                            70
                                                 75
     59 Leu Gln Trp Ser Ser Leu Lys Ala Ser Asp Thr Gly Met Tyr Tyr Cys
                                             90
                        85
     63 Ala Arg Leu Glu Pro Gly Tyr Ser Ser Thr Trp Ser Val Asn Trp Gly
                    100
                                         105
     67 Gln Gly Thr Leu Val Thr Val Ser Ser
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                                    120
     71 <210> SEQ ID NO: 2
    72 <211> LENGTH: 5
     73 <212> TYPE: PRT
     74 <213> ORGANISM: Homo sapiens
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76 <400> SEQUENCE: 2

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Input Set : N:\efs\10537280_efs\URQUP16_seq.txt
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78 Ser Tyr Trp Ile Asn
79 1
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83 <211> LENGTH: 17
84 <212> TYPE: PRT
85 <213> ORGANISM: Homo sapiens
87 <400> SEQUENCE: 3
89 Arg Ile Asp Pro Thr Asp Ser Tyr Thr Asn Tyr Ser Pro Ser Phe Lys
90 1
93 Gly
97 <210> SEQ ID NO: 4
98 <211> LENGTH: 12
99 <212> TYPE: PRT
100 <213> ORGANISM: Homo sapiens
102 <400> SEQUENCE: 4
104 Leu Glu Pro Gly Tyr Ser Ser Thr Trp Ser Val Asn
105 1
108 <210> SEQ ID NO: 5
109 <211> LENGTH: 131
110 <212> TYPE: PRT
111 <213> ORGANISM: Homo sapiens
113 <400> SEQUENCE: 5
115 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu
119 Ser Leu Lys Ile Ser Cys Arg Gly Ser Gly Tyr Arg Phe Thr Ser Tyr
123 Trp Ile Asn Trp Val Arg Gln Leu Pro Gly Lys Gly Leu Glu Trp Met
                                40
127 Gly Arg Ile Asp Pro Thr Asp Ser Tyr Thr Asn Tyr Ser Pro Ser Phe
131 Lys Gly His Val Thr Val Ser Ala Asp Lys Ser Ile Asn Thr Ala Tyr
                        70
                                            75
135 Leu Gln Trp Ser Ser Leu Lys Ala Ser Asp Thr Gly Met Tyr Tyr Cys
139 Ala Arg Leu Glu Pro Gly Tyr Ser Ser Thr Trp Ser Val Asn Trp Gly
                100
                                    105
143 Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser
144
          115
147 Val Phe Pro
148
        130
151 <210> SEQ ID NO: 6
152 <211> LENGTH: 111
153 <212> TYPE: PRT
154 <213> ORGANISM: Homo sapiens
156 <400> SEQUENCE: 6
158 Leu Thr Val Leu Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Arg Gln
                                        10
162 Arg Val Thr Ile Ser Cys Ser Gly Asn Ser Ser Asn Ile Gly Asn Asn
163
                20
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Input Set : N:\efs\10537280_efs\URQUP16_seq.txt
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166 Ala Val Asn Trp Tyr Gln Gln Leu Pro Gly Lys Ala Pro Lys Leu Leu
170 Ile Tyr Tyr Asp Asp Gln Leu Pro Ser Gly Val Ser Asp Arg Phe Ser
171
                            55
174 Gly Ser Arg Ser Gly Thr Ser Ala Ser Leu Ala Ile Arg Gly Leu Gln
178 Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Thr Ser Trp Asp Asp Ser Leu
179
182 Asp Ser Gln Leu Phe Gly Gly Gly Thr Arg Leu Thr Val Leu Gly
183
186 <210> SEQ ID NO: 7
187 <211> LENGTH: 13
188 <212> TYPE: PRT
189 <213> ORGANISM: Homo sapiens
191 <400> SEQUENCE: 7
193 Ser Gly Asn Ser Ser Asn Ile Gly Asn Asn Ala Val Asn
194 1
197 <210> SEQ ID NO: 8
198 <211> LENGTH: 7
199 <212> TYPE: PRT
200 <213> ORGANISM: Homo sapiens
202 <400> SEQUENCE: 8
204 Tyr Asp Asp Gln Leu Pro Ser
205 1
208 <210> SEQ ID NO: 9
209 <211> LENGTH: 11
210 <212> TYPE: PRT
211 <213> ORGANISM: Homo sapiens
213 <400> SEQUENCE: 9
215 Thr Ser Trp Asp Asp Ser Leu Asp Ser Gln Leu
216 1
219 <210> SEQ ID NO: 10
220 <211> LENGTH: 363
221 <212> TYPE: DNA
222 <213> ORGANISM: Homo sapiens
224 <400> SEQUENCE: 10
225 caaatgcagc tggtgcagtc tggagcagag gtgaaaaagc ccggggagtc tctgaagatc
                                                                           60
227 teetgtaggg gttetggata eaggtttace agetactgga teaactgggt gegeeagetg
                                                                          120
229 cccgggaaag gcctagagtg gatgggcagg attgatccta ctgactctta taccaactac
                                                                          180
231 agtocatect teaaaggeea egteacegte teagetgaca agteeateaa eactgeetae
                                                                          240
233 ctgcagtgga gcagcctgaa ggcctcggac accggcatqt attactqtqc qaqqctcqaa
                                                                          300
235 ccgggctata gcagcacctg gtccgtaaat tggggccagg gaaccctggt caccgtctcc
                                                                          360
237 tca
                                                                          363
240 <210> SEQ ID NO: 11
241 <211> LENGTH: 15
242 <212> TYPE: DNA
243 <213> ORGANISM: Homo sapiens
245 <400> SEQUENCE: 11
246 agctactgga tcaac
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Input Set : N:\efs\10537280 efs\URQUP16 seq.txt

Output Set: N:\CRF4\01092006\J537280.raw

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249 <210> SEQ ID NO: 12
250 <211> LENGTH: 51
251 <212> TYPE: DNA
252 <213> ORGANISM: Homo sapiens
254 <400> SEQUENCE: 12
255 aggattgatc ctactgactc ttataccaac tacagtccat ccttcaaagg c
                                                                           51
258 <210> SEQ ID NO: 13
259 <211> LENGTH: 36
260 <212> TYPE: DNA
261 <213> ORGANISM: Homo sapiens
263 <400> SEQUENCE: 13
264 ctcgaaccgg gctatagcag cacctggtcc gtaaat
                                                                           36
267 <210> SEQ ID NO: 14
268 <211> LENGTH: 394
269 <212> TYPE: DNA
270 <213> ORGANISM: Homo sapiens
272 <400> SEQUENCE: 14
273 caaatgcagc tggtgcagtc tggagcagag gtgaaaaagc ccggggagtc tctgaagatc
                                                                           60
275 tcctgtaggg gttctggata caggtttacc agctactgga tcaactgggt gcgccagctg
                                                                          120
277 cccgggaaag gcctagagtg gatgggcagg attgatccta ctgactctta taccaactac
279 agtocatoot toaaaggooa ogtoacogto toagotgaca agtocatoaa cactgootac
                                                                          240
281 ctgcagtgga gcagcctgaa ggcctcggac accggcatgt attactgtgc gaggctcgaa
                                                                          300
283 ccgggctata gcagcacctg gtccgtaaat tggggccagg gaaccctggt caccgtctcc
                                                                          360
285 tcagcctcca ccaagggccc atcggtcttc cccc
                                                                          394
288 <210> SEQ ID NO: 15
289 <211> LENGTH: 333
290 <212> TYPE: DNA
291 <213> ORGANISM: Homo sapiens
293 <400> SEQUENCE: 15
294 ctgcctgtgc tgactcagcc accctcggtg tctggagccc ccaggcagag ggtcaccatc
                                                                           60
296 teetgttetg gaaacagete caacategga aataatgetg taaaetggta eeageagete
                                                                          120
298 ccaggaaagg ctcccaaact cctcatttat tatgatgatc aactgccctc aggggtctct
                                                                          180
300 gaccgattet etggetecag gtetggeace teegeeteee tggecateeg tgggetecag
                                                                          240
302 totgaggatg aggotgatta ttactgtaca toatgggatg acagootgga tagtoaactg
                                                                          300
304 ttcggcggag ggaccaggct gaccgtccta ggt
                                                                          333
307 <210> SEQ ID NO: 16
308 <211> LENGTH: 39
309 <212> TYPE: DNA
310 <213> ORGANISM: Homo sapiens
312 <400> SEQUENCE: 16
313 totggaaaca gotocaacat oggaaataat gotgtaaac
                                                                           39
316 <210> SEQ ID NO: 17
317 <211> LENGTH: 21
318 <212> TYPE: DNA
319 <213> ORGANISM: Homo sapiens
321 <400> SEQUENCE: 17
322 tatgatgatc aactgccctc a
                                                                           21
325 <210> SEQ ID NO: 18
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326 <211> LENGTH: 33

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Input Set : N:\efs\10537280_efs\URQUP16_seq.txt
Output Set: N:\CRF4\01092006\J537280.raw

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327 <212> TYPE: DNA
328 <213> ORGANISM: Homo sapiens
330 <400> SEQUENCE: 18
331 acatcatggg atgacagect ggatagteaa etg
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334 <210> SEQ ID NO: 19
335 <211> LENGTH: 119
336 <212> TYPE: PRT
337 <213> ORGANISM: Mus sp.
339 <400> SEQUENCE: 19
341 Asp Val Gln Ile Gln Gln Pro Gly Thr Glu Leu Val Lys Pro Gly Ala
345 Ser Val Arg Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Tyr
                                     25
349 Trp Met His Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
350
            35
353 Gly Glu Ile Asp Pro Ser Asp Ser Tyr Thr Asn Tyr Asn Gln Lys Phe
                            55
357 Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr
                        70
                                             75
361 Met His Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
365 Ser Arg Asn Tyr Gly Ser Gly Tyr Tyr Phe Asp Tyr Trp Gly Gln Gly
366
                100
                                     105
                                                         110
369 Thr Thr Leu Thr Val Ser Ser
370
            115
373 <210> SEQ ID NO: 20
374 <211> LENGTH: 5
375 <212> TYPE: PRT
376 <213> ORGANISM: Mus sp.
378 <400> SEQUENCE: 20
380 Thr Tyr Trp Met His
381 1
384 <210> SEQ ID NO: 21
385 <211> LENGTH: 17
386 <212> TYPE: PRT
387 <213> ORGANISM: Mus sp.
389 <400> SEQUENCE: 21
391 Glu Ile Asp Pro Ser Asp Ser Tyr Thr Asn Tyr Asn Gln Lys Phe Lys
392 1
                                         10
395 Gly
399 <210> SEQ ID NO: 22
400 <211> LENGTH: 10
401 <212> TYPE: PRT
402 <213> ORGANISM: Mus sp.
404 <400> SEQUENCE: 22
406 Asn Tyr Gly Ser Gly Tyr Tyr Phe Asp Tyr
407 1
410 <210> SEQ ID NO: 23
411 <211> LENGTH: 124
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VERIFICATION SUMMARYDATE: 01/09/2006PATENT APPLICATION: US/10/537,280TIME: 08:29:27

Input Set : N:\efs\10537280_efs\URQUP16_seq.txt
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L:12 M:270 C: Current Application Number differs, Replaced Current Application Number L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date